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Cys His Gln Cys Asp Pro Asp Lys Tyr Ser Glu Lys Gly Ser Ser Ser	
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gac cga gat tca gga acc tgc cac tcc tgc ccc cct aac aca att ctg 1158  Asp Arg Asp Ser Gly Thr Cys His Ser Cys Pro Pro Asn Thr Ile Leu  365 370 375
aaa gcc cac cag cct tat ggt gtc cag gcc tgt gtg ccc tgt ggt cca 1206 Lys Ala His Gln Pro Tyr Gly Val Gln Ala Cys Val Pro Cys Gly Pro 380 385 390
ggg acc aag aac aac aag atc cac tct ctg tgc tac aat gat tgc acc 1254 Gly Thr Lys Asn Asn Lys Ile His Ser Leu Cys Tyr Asn Asp Cys Thr 395 400 405
ttc tca cgc aac act cca acc agg act ttc aac tac aac ttc tcc gct 1302  Phe Ser Arg Asn Thr Pro Thr Arg Thr Phe Asn Tyr Asn Phe Ser Ala  410  415
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Tyr Ser Glu Lys Gly Ser Ser Ser Cys Asn Val Arg Pro Ala Cys Thr 65 70 75 80

Asp Lys Asp Tyr Phe Tyr Thr His Thr Ala Cys Asp Ala Asn Gly Glu 85 90 95

Thr Gln Leu Met Tyr Lys Trp Ala Lys Pro Lys Ile Cys Ser Glu Asp 100 105 110

Leu Glu Gly Ala Val Lys Leu Pro Ala Ser Gly Val Lys Thr His Cys 115 120 125

Pro Pro Cys Asn Pro Gly Phe Phe Lys Thr Asn Asn Ser Thr Cys Gln 130

Pro Cys Pro Tyr Gly Ser Tyr Ser Asn Gly Ser Asp Cys Thr Arg Cys 145 150 155

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Thr Leu Pro Thr Asn Met Glu Thr Thr Val Leu Ser Gly Ile Asn Phe 180 185 190

Glu Tyr Lys Gly Met Thr Gly Trp Glu Val Ala Gly Asp His Ile Tyr 195 200 205

Thr Ala Ala Gly Ala Ser Asp Asn Asp Phe Met Ile Leu Thr Leu Val 210 215 220

Val Pro Gly Phe Arg Pro Pro Gln Ser Val Met Ala Asp Thr Glu Asn 225 230 235 240

Lys Glu Val Ala Arg Ile Thr Phe Val Phe Glu Thr Leu Cys Ser Val 245 250 255 Asn Cys Glu Leu Tyr Phe Met Val Gly Val Asn Ser Arg Thr Asn Thr 260 265 270

Pro Val Glu Thr Trp Lys Gly Ser Lys Gly Lys Gln Ser Tyr Thr Tyr 275

Ile Ile Glu Glu Asn Thr Thr Thr Ser Phe Thr Trp Ala Phe Gln Arg 290 295 300

Thr Thr Phe His Glu Ala Ser Arg Lys Tyr Thr Asn Asp Val Ala Lys 305 310 315

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Thr Leu Ser Leu Cys Gly Asn Gln Gly Arg Lys Met Ser Val Cys Thr 450 455 460

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Ser Pro Ala Glu Leu Phe His Leu Glu Ser Leu Gly Ile Pro Asp Val 530 535

Ile Phe Phe Tyr Arg Ser Asn Asp Val Thr Gln Ser Cys Ser Ser Gly 545 550 560

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Ser Val Ala Asp Tyr His Ala Ile Val Ser Ser Cys Val Ala Gly Ile

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Gly Asn Gly Met Val Ser Arg Cys Ser Arg Ser Gln Asn Thr Val Cys

Arg Pro Cys Gly Pro Gly Phe Tyr Asn Asp Val Val Ser Ser Lys Pro

Cys Lys Pro Cys Thr Trp Cys Asn Leu Arg Ser Gly Ser Glu Arg Lys

Gln Leu Cys Thr Ala Thr Gln Asp Thr Val Cys Arg Cys Arg Ala Gly

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Pro Pro Gly His Phe Ser Pro Gly Asp Asn Gln Ala Cys Lys Pro Trp 130 135 140

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220 225

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<sup>&</sup>lt;211> 226

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapiens

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Tr	p Al	a Al		a Hi	s Al	a Le	u Pr	o Al 2	a Gl :5	n Va	l Al	a Ph	e Th	r Pr O	o Tyr
Al	a Pr		lu Pr	o Gl	y Se	er Th	ır Cy 4	rs Ar	g Le	eu Ar	g Gl	u Ty 4	r Ту 5	r As	p Glr

Thr Ala Gln Met Cys Cys Ser Lys Cys Ser Pro Gly Gln His Ala Lys

Val Phe Cys Thr Lys Thr Ser Asp Thr Val Cys Asp Ser Cys Glu Asp 65 70 75 80

55

Ser Thr Tyr Thr Gln Leu Trp Asn Trp Val Pro Glu Cys Leu Ser Cys 85 90 95

Gly Ser Arg Cys Ser Ser Asp Gln Val Glu Thr Gln Ala Cys Thr Arg 100 105 110

Glu Gln Asn Arg Ile Cys Thr Cys Arg Pro Gly Trp Tyr Cys Ala Leu 115 120 125.

Ser Lys Gln Glu Gly Cys Arg Leu Cys Ala Pro Leu Arg Lys Cys Arg 130 135 140

Pro Gly Phe Gly Val Ala Arg Pro Gly Thr Glu Thr Ser Asp Val Val 145 150 150

Cys Lys Pro Cys Ala Pro Gly Thr Phe Ser Asn Thr Thr Ser Ser Thr 165 170 175

Asp Ile Cys Arg Pro His Gln Ile Cys Asn Val Val Ala Ile Pro Gly 180 185 190

Asn Ala Ser Arg Asp Ala Val Cys Thr Ser Thr Ser Pro Thr Arg Ser 195 200 205

Met Ala Pro Gly Ala Val His Leu Pro Gln Pro Val Ser Thr Arg Ser 210 215 220

Gln His Thr Gln Pro Thr Pro Glu Pro Ser Thr Ala Pro Ser Thr Ser 225 230 235 235

Phe Leu Leu Pro Met Gly Pro Ser Pro Pro Ala Glu Gly Ser Thr Gly 245 250 255

Asp Phe Ala Leu Pro Val Gly Leu Ile Val Gly Val Thr Ala Leu Gly 260 265 270

Leu Leu Ile Ile Gly Val Val Asn Cys Val Ile Met Thr Gln Val Lys 275 280 285

Lys Lys Pro Leu Cys Leu Gln Arg Glu Ala Lys Val Pro His Leu Pro 290 295 300

Ala Asp Lys Ala Arg Gly Thr Gln Gly Pro Glu Gln Gln His Leu Leu 305 310 315 320

Ile Thr Ala Pro Ser Ser Ser Ser Ser Leu Glu Ser Ser Ala Ser 325 330 335

Ala Leu Asp Arg Arg Ala Pro Thr Arg Asn Gln Pro Gln Ala Pro Gly 340 345 350

Val Glu Ala Ser Gly Ala Gly Glu Ala Arg Ala Ser Thr Gly Ser Ser 355 360 365

Asp Ser Ser Pro Gly Gly His Gly Thr Gln Val Asn Val Thr Cys Ile 370 375 380

Val Asn Val Cys Ser Ser Ser Asp His Ser Ser Gln Cys Ser Ser Gln 395 400

Ala Ser Ser Thr Met Gly Asp Thr Asp Ser Ser Pro Ser Glu Ser Pro

Lys Asp Glu Gln Val Pro Phe Ser Lys Glu Glu Cys Ala Phe Arg Ser 425

Gln Leu Glu Thr Pro Glu Thr Leu Leu Gly Ser Thr Glu Glu Lys Pro 440

Leu Pro Leu Gly Val Pro Asp Ala Gly Met Lys Pro Ser 455

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Val Leu Asp Arg Ser Tyr Pro Arg Ile Val Val Met Glu Arg Val Glu

Met Pro Thr Ala Gln Pro Ala Leu Leu Ala Val Gln Lys Gln Leu Gly

Pro Pro Gln Met Cys Arg Val Ala Cys Thr Cys Ala Val Ile Asn Arg

Val Gln Lys Val Asn Cys Thr Pro Thr Ser Asn Ala Val Cys Gly Asp

Cys Leu Pro Arg Phe Tyr Arg Lys Thr Arg Ile Gly Gly Leu Gln Asp

Gln Glu Cys Ile Pro Cys Thr Lys Gln Thr Pro Thr Ser Glu Val Gln

Cys Ala Phe Gln Leu Ser Leu Val Glu Ala Asp Ala Pro Thr Val Pro

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 ntacaaatgg tggaacacgc tgcccacaaa catggaaacg accgttctca gtgggatcaa 240
 cttcgagtac aagggcatga caggctggga ggtggntggt gntcacattt acacagctgc 300
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  ggaagtacac caatgacgtt gccaagatct actccatcaa tgtcaccaat gttatgaatg 180
  gegtggeete etaetgeegt eeetgtgeee tagaageete tgatgtggge teeteetgea 240
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 atattgaccg agattcagga acctgccact cctgcccccc taacacaatt ntgaaagccc 180
 accagcetta tggtgtccag geetgtntge eetgtggtee agggaccaag aacaacaaga 240
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  ctgtgacctg ccagcagctg acagctcgcc atcatggaag gcgaggntgt agaggacgac 120
  ctcatcttta ccagcaagaa gtcactcttt gggaagatca aatcatttac ctccaagagg 180
  actcctgatg gatttnactc agtgccgctg aagacatcct caggaggccc agacatggac 240
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  ctgtgagagg cactgcctgc ctcacctgct tcct
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gccctgccca tatggttcct actccaatgg ctcagactgt acccgctgcc ctgcagggac 180
tgaacctgct gtgggatttg aatacaaatg gtggaacacg ctgcccacaa acatgggaaa 240
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cgacc
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  tgtgagaggc actgcctgcc tcacctgcct cctcaccttg catagcacct ttgcaagcct 120
  gcgggaattt gggtgccagc atcctgcaac acccactgct gggaaatctc ttcattgtgg 180
  ccttatcaga tgtttgaatt tnagatcttt ttttatagag tacccaaacc ctcctttctg 240
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tgctcagatg ggacctgtga tggctgcaac ttccacttcc tgtnggagag cgcggctgct 180
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ctttgagacc ctctgttctg tgaactgtga gctctacttc atggtgggtg tggaattcta 180
 gggaccaaca cttcctgtgg aggacgtggg aaaggttcca aagggcaaac agtnccttat 240
 tacctgacat gcattgaggn aggaacantt nnccnggagg tttcaactgg ggcctttccc 300
 gaggnacnac ttttttcatg gagggccaag ncaggggagt tacaacccat tgnacgttng 360
 gccaaggntc tnatttccat ncaatgtnca accaatgntn atggaanggg tgttggggcc 420
                                                                     427
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  <222> (80)
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                                                                     70
 ctcatntaca
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 taaagtattt cgggtaaagg tgaagtgaag gattttcgtc tttataattt ctgttcggcc 180
 atggcaaata ccatagttga gtatttgctt caggagagtt ctttttacag ttttactttt 240
 caatgctgag gcatatttct ttgagcactg tgcttttatg tgtctttcta caaaggggtt 300
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 aaacagcagt taagttgtct ctaaatgaac tagcanaaaa aaaaaatgta gtttttgttt 420
 gtaaggaagg ggaggtattt cctgagaatg aattttttt ttttnggaaa cnggtttctn 480
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taaagtattt cgggtaaagg tgaagtgaag gattttcgtc tttataattt ctgttcggcc 180
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 aaacagcagt taagttgtct ctaaatgaac tagcanaaaa aaaaaangta gtttttgttt 420
 gtaaggaagg ggaggtattt cctgagaatg aattttttt tttttggata acnggttttc 480
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                                                                    554
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  cetttgatte tttetcaatt gtntttttge etttagetee caectataca teteatgete 120
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310

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 ccccaacnnc tgnaggttcc aatgtggcct tnccatttgg aagcttantg ggaaggcaga 480
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  gcacagtnac ctctnatgct gtctgtgggg ganggtttgc ccaagtttct aaccgaaaga 360
  cacgccattg gaaggetgee aggaccaagg atggcatece gtggcacaaa gncagaceee 420
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  <211> 24
   <212> DNA
   <213> Homo sapiens
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Ser Leu Pro Asp Pro Val Lys Gly Thr Glu Cys Ser Phe Ser Cys Asn 85 90 95

Ala Gly Glu Phe Leu Asp Met Lys Asp Gln Ser Cys Lys Pro Cys Ala

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Glu Leu Pro His Gly Phe Ala Ser Leu Ser Ala Asn Met Glu Leu Asp 130 135 140

Asp Ser Ala Ala Glu Ser Thr Gly Asn Cys Thr Ser Ser Lys Trp Val 145 150 155 160

Pro Arg Gly Asp Tyr Ile Ala Phe Asn Thr Asp Glu Cys Thr Ala Thr 165

Leu Met Tyr Ala Val Asn Leu Lys Gln Ser Gly Thr Val Asn Phe Glu 180 185 190

Tyr Tyr Tyr Pro Asp Ser Ser Ile Ile Phe Glu Phe Phe Val Gln Asn 195 200 205

Asp Gln Cys Gln Pro Asn Ala Asp Asp Ser Arg Trp Met Lys Thr Thr 210 215 220

Glu Lys Gly Trp Glu Phe His Ser Val Glu Leu Asn Arg Gly Asn Asn 225 230 235

Val Leu Tyr Trp Arg Thr Thr Ala Phe Ser Val Trp Thr Lys Val Pro

Lys Pro Val Leu Val Arg Asn Ile Ala Ile Thr Gly Val Ala Tyr Thr 260 265 270

Ser Glu Cys Phe Pro Cys Lys Pro Gly Thr Tyr Ala Asp Lys Gln Gly 275 280 285

Ser Ser Phe Cys Lys Leu Cys Pro Ala Asn Ser Tyr Ser Asn Lys Gly 290 295

Glu Thr Ser Cys His Gln Cys Asp Pro Asp Lys Tyr Ser Glu Lys Gly 305 310 315

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Tyr Thr His Thr Ala Cys Asp Ala Asn Gly Glu Thr Gln Leu Met Tyr 340 345 350

Lys Trp Ala Lys Pro Lys Ile Cys Ser Glu Asp Leu Glu Gly Ala Val 355 360 365

Lys Leu Pro Ala Ser Gly Val Lys Thr His Cys Pro Pro Cys Asn Pro 370

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Ser Tyr Ser Asn Gly Ser Asp Cys Thr Arg Cys Pro Ala Gly Thr Glu 405

Pro Ala Val Gly Phe Glu Tyr Lys Trp Trp Asn Thr Leu Pro Thr Asn 420 425 430

Met Glu Thr Thr Val Leu Ser Gly Ile Asn Phe Glu Tyr Lys Gly Met 435

Thr Gly Trp Glu Val Ala Gly Asp His Ile Tyr Thr Ala Ala Gly Ala 450 455

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Pro Pro Gln Ser Val Met Ala Asp Thr Glu Asn Lys Glu Val Ala Arg 495

Ile Thr Phe Val Phe Glu Thr Leu Cys Ser Val Asn Cys Glu Leu Tyr 500 505 510

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Lys Gly Ser Lys Gly Lys Gln Ser Tyr Thr Tyr Ile Ile Glu Glu Asn 530

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Val Thr Asn Val Met Asn Gly Val Ala Ser Tyr Cys Arg Pro Cys Ala 580 585 590

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Tyr Tyr Ile Asp Arg Asp Ser Gly Thr Cys His Ser Cys Pro Pro Asn 610 615

Thr Ile Leu Lys Ala His Gln Pro Tyr Gly Val Gln Ala Cys Val Pro 625 630 635

Cys Gly Pro Gly Thr Lys Asn Asn Lys Ile His Ser Leu Cys Tyr Asn 655

Asp Cys Thr Phe Ser Arg Asn Thr Pro Thr Arg Thr Phe Asn Tyr Asn 660 665 670

Phe Ser Ala Leu Ala Asn Thr Val Thr Leu Ala Gly Gly Pro Ser Phe 675 680 685

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Pro Gly Thr Tyr Ala Ser Arg Leu Cys Asp Ser Lys Thr Asn Thr Asn

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His Leu Pro Ala Cys Leu Ser Cys Asn Gly Arg Cys Asp Ser Asn Gln

Val Glu Thr Arg Ser Cys Asn Thr Thr His Asn Arg Ile Cys Asp Cys 105

Ala Pro Gly Tyr Tyr Cys Leu Leu Lys Gly Ser Gly Cys Lys Ala Cys

Val Ser Gln Thr Lys Cys Gly Ile Gly Tyr Gly Val Ser Gly His Thr

Pro Thr Gly Asp Val Ile Cys Ser Pro Cys Gly Leu Gly Thr Tyr Ser

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<223> n equals a, t, g or c

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cagcctgcct gaccccgtca agggcaccga gtgctcette tcctgcaacg ccgggagtt 180

tctggatatg aaggaccagt catgtaagce atgcgctgag ggccgctact ccctcggcac 240

aggcattcgg tttgatgagt gggatgagct tgcccatgg ctttgcagc tttt 294
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291

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<212> DNA
<213> Homo sapiens
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<221> misc_feature
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<223> n equals a,t,g, or c
<220>
<221> misc_feature
<222> (171)
<223> n equals a,t,g, or c
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 atattgaccg agattcagga acctgccact cctgcccccc taacacaatt ntgaaagccc 180
 accagcetta tggtgtccag geetgtntge cetgtggtee agggaccaag aacaacaaga 240
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  <223> n equals a, t, g or c
  <220>
  <221> misc_feature
  <222> (196)
  <223> n equals a, t, g or c
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  ctgtgacctg ccagcagctg acagctcgcc atcatggaag gcgaggntgt agaggacgac 120
  ctcatcttta ccagcaagaa gtcactcttt gggaagatca aatcatttac ctccaagagg 180
  actcctgatg gatttnactc agtgccgctg aagacatcct caggaggccc agacatggac 240
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  ctgtgagagg cactgcctgc ctcacctgct tcct
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<210> 53

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<211> 245
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gccctgccca tatggttcct actccaatgg ctcagactgt acccgctgcc ctgcagggac 180
tgaacctgct gtgggatttg aatacaaatg gtggaacacg ctgcccacaa acatgggaaa 240
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 cgacc
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 <211> 292
 <212> DNA
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 <223> n equals a, t, g or c
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 <222> (202)
 <223> n equals a, t, g or c
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  <221> misc_feature
  <222> (245)
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  <220>
  <221> misc_feature
  <222> (246)
  <223> n equals a, t, g or c
  <220>
  <221> misc_feature
  <222> (291)
  <223> n equals a, t, g or c
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   <221> misc_difference
   <222> (292)
   <223> n equals a, t, g or c
   ggcanaggga atttgactca gtgccgctga agacatcctc aggaggccca gacatggacc 60
   tgtgagaggc actgcctgcc tcacctgcct cctcaccttg catagcacct ttgcaagcct 120
   gcgggaattt gggtgccagc atcctgcaac acccactgct gggaaatctc ttcattgtgg 180
   ccttatcaga tgtttgaatt tnagatcttt ttttatagag tacccaaacc ctcctttctg 240
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220

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<212> DNA
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tgcccgctct gctcagtggc tgactaccat gctatcgtca
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  <222> (234)
  <223> n equals a, t, g or c
  <220>
  <221> misc_feature
  <222> (260)
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  <221> misc_feature
  <222> (268)
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  <220>
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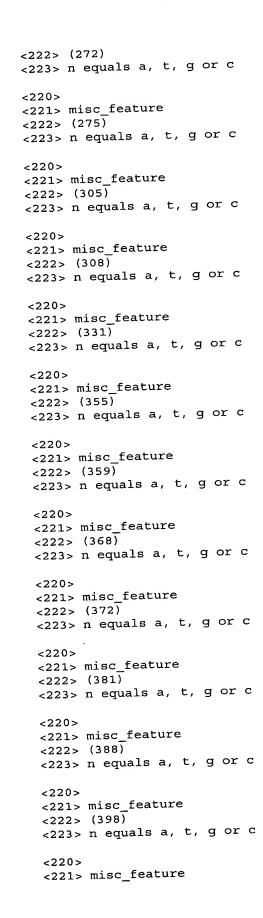
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<221> misc\_feature

<223> n equals a, t, g or c

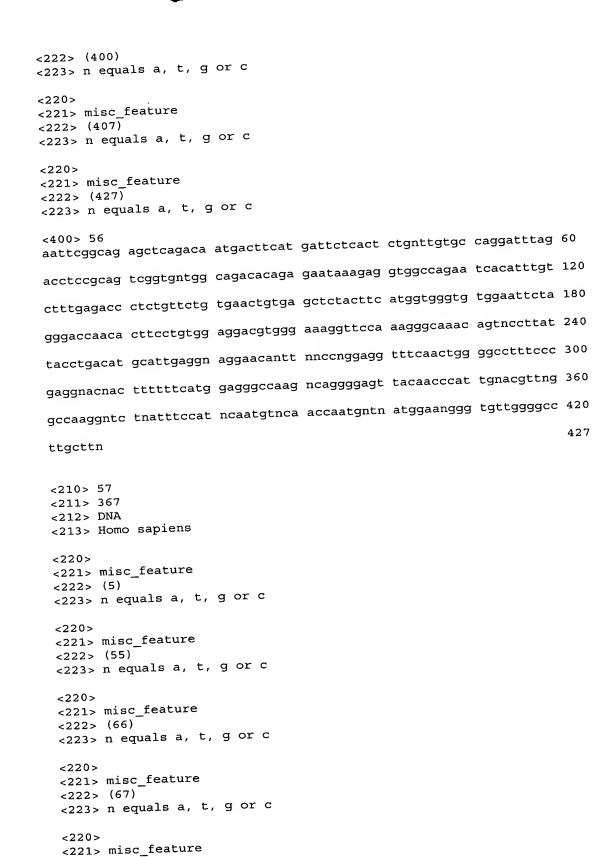
<222> (271)

<220>



<222> (116)

<223> n equals a, t, g or c



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<223> n equals a, t, g or c
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 agnotggtca cogttgaact toggaatact actatocaga otocatoato atotttgaag 180
 tttttcgttc agaatgacca gtgccagccc aatgcagatg actccaggtg gatgaagacc 240
 acagagaaag gatgggaatt ccacagtgtg agctnaaatc gaggcaataa tgtccgttat 300
  tgggggaacc acagnettet teaatgatgg gaccaaagtn acceaagnet gtgetnggtg 360
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  <210> 58
  <211> 333
  <212> DNA
  <213> Homo sapiens
  <220>
  <221> misc_feature
  <222> (20)
   <223> n equals a, t, g or c
   <220>
   <221> misc_feature
   <222> (23)
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<223> n equals a, t, g or c
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<221> misc_feature
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 <223> n equals a, t, g or c
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 <222> (152)
 <223> n equals a, t, g or c
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 <223> n equals a, t, g or c
  <220>
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  <223> n equals a, t, g or c
  <220>
  <221> misc_feature
  <222> (260)
  <223> n equals a, t, g or c
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  <222> (269)
  <223> n equals a, t, g or c
  <220>
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  <223> n equals a, t, g or c
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<221> misc_feature
<222> (307)
<223> n equals a, t, g or c
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<222> (320)
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aggttccaaa ggcaaacagt cctataccta catcattgaa ggaggaacac taccacgagg 240
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 <211> 70
 <212> DNA
 <213> Homo sapiens
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 <221> misc_feature
 <222> (40)
 <223> n equals a, t, g or c
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 <221> misc_feature
 <222> (60)
 <223> n equals a, t, g or c
 <220>
 <221> misc feature
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  <211> 3152
  <212> DNA
  <213> Homo sapiens
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  cccaccatgg attgccaaga aaatgagtac tgggaccaat ggggacggtg tgtcacctgc 120
  caacggtgtg gtcctggaca ggagctatcc aaggattgtg gttatggaga gggtggagat 180
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<210> 61
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<sup>&</sup>lt;211> 231

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapiens

<sup>&</sup>lt;400> 61 Met Asp Cys Gln Glu Asn Glu Tyr Trp Asp Gln Trp Gly Arg Cys Val 10





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- Tyr Gly Glu Gly Gly Asp Ala Tyr Trp His Ser Leu Pro Ser Ser Gln
  35 40 45
- Tyr Lys Ser Ser Trp Gly His His Lys Cys Gln Ser Cys Ile Thr Cys
  50 60
- Ala Val Ile Asn Arg Val Gln Lys Val Asn Cys Thr Pro Thr Ser Asn 65 70 75 80
- Ala Val Cys Gly Asp Cys Leu Pro Arg Phe Tyr Arg Lys Thr Arg Ile 85 90 95
- Gly Gly Leu Gln Asp Gln Glu Cys Ile Pro Cys Thr Lys Gln Thr Pro 100 105 110
- Thr Ser Glu Val Gln Cys Ala Phe Gln Leu Ser Leu Val Glu Ala Asp 115 120 125
- Ala Pro Thr Val Pro Pro Gln Glu Ala Thr Leu Val Ala Leu Val Ser 130 135 140
- Ser Leu Leu Val Val Phe Thr Leu Ala Phe Leu Gly Leu Phe Phe Leu 145 150 155 160
- Tyr Cys Lys Gln Phe Phe Asn Arg His Cys Gln Arg Gly Gly Leu Leu 165 170 175
- Gln Phe Glu Ala Asp Lys Thr Ala Lys Glu Glu Ser Leu Phe Pro Val 180 185 190
- Pro Pro Ser Lys Glu Thr Ser Ala Glu Ser Gln Val Ser Trp Ala Pro 195 200 205
- Gly Ser Leu Ala Gln Leu Phe Ser Leu Asp Ser Val Pro Ile Pro Gln 210 220
- Gln Gln Gln Gly Pro Glu Met 225 230